

#15

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/245,025

DATE: 08/13/2001

TIME: 11:46:15

Input Set : N:\Cr3\RULE60\09245025.txt

Output Set: N:\CRF3\08132001\I245025.raw

SEQUENCE LISTING

4 (1) GENERAL INFORMATION:

6 (i) APPLICANT: Gerard, Gary F.

7 Smith, Michael D.

8 Chatterjee, Deb K.

10 (ii) TITLE OF INVENTION: Compositions and Methods for Reverse

11 Transcription of Nucleic Acid Molecules

13 (iii) NUMBER OF SEQUENCES: 23

15 (iv) CORRESPONDENCE ADDRESS:

16 (A) ADDRESSEE: STERNE, KESSLER, GOLDSTEIN & FOX P.L.L.C.

17 (B) STREET: 1100 New York Avenue, Suite 600

18 (C) CITY: Washington

19 (D) STATE: DC

20 (E) COUNTRY: USA

21 (F) ZIP: 20005

23 (v) COMPUTER READABLE FORM:

24 (A) MEDIUM TYPE: Floppy disk

25 (B) COMPUTER: IBM PC compatible

26 (C) OPERATING SYSTEM: PC-DOS/MS-DOS

27 (D) SOFTWARE: PatentIn Release #1.0, Version #1.30

29 (vi) CURRENT APPLICATION DATA:

C--> 30 (A) APPLICATION NUMBER: US/09/245,025

C--> 31 (B) FILING DATE: 25-Feb-1999

37 (C) CLASSIFICATION:

39 (vii) PRIOR APPLICATION DATA:

35 (A) APPLICATION NUMBER: 09/064,057

36 (B) FILING DATE:

40 (A) APPLICATION NUMBER: US 60/044,589

41 (B) FILING DATE: 22-APR-1997

43 (viii) ATTORNEY/AGENT INFORMATION:

44 (A) NAME: Esmond, Robert W.

45 (B) REGISTRATION NUMBER: 32,893

46 (C) REFERENCE/DOCKET NUMBER: 0942.4330002

48 (ix) TELECOMMUNICATION INFORMATION:

49 (A) TELEPHONE: 202-371-2600

50 (B) TELEFAX: 202-371-2540

53 (2) INFORMATION FOR SEQ ID NO: 1:

55 (i) SEQUENCE CHARACTERISTICS:

56 (A) LENGTH: 35 base pairs

57 (B) TYPE: nucleic acid

58 (C) STRANDEDNESS: both

59 (D) TOPOLOGY: both

61 (ii) MOLECULE TYPE: cDNA

66 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

68 AGGAGACCCA TATGACTGTT GCGCTACATC TGGCT

70 (2) INFORMATION FOR SEQ ID NO: 2:

72 (i) SEQUENCE CHARACTERISTICS:

ENTERED

35

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73          (A) LENGTH: 34 base pairs
74          (B) TYPE: nucleic acid
75          (C) STRANDEDNESS: both
76          (D) TOPOLOGY: both
78      (ii) MOLECULE TYPE: cDNA
83      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
85 AACGCGACAG GTTAACAGCG CGCAAATCAT GCAG      34
87 (2) INFORMATION FOR SEQ ID NO: 3:
89      (i) SEQUENCE CHARACTERISTICS:
90          (A) LENGTH: 32 base pairs
91          (B) TYPE: nucleic acid
92          (C) STRANDEDNESS: both
93          (D) TOPOLOGY: both
95      (ii) MOLECULE TYPE: cDNA
100     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
102 CACACACAGG TACCCTCTCG AAAAGTTAAA CC      32
104 (2) INFORMATION FOR SEQ ID NO: 4:
106     (i) SEQUENCE CHARACTERISTICS:
107         (A) LENGTH: 41 base pairs
108         (B) TYPE: nucleic acid
109         (C) STRANDEDNESS: both
110         (D) TOPOLOGY: both
112     (ii) MOLECULE TYPE: cDNA
117     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
119 CACACACACT CGAGTTATGC AAAAAGAGGG CTCGCCTCAT C      41
121 (2) INFORMATION FOR SEQ ID NO: 5:
123     (i) SEQUENCE CHARACTERISTICS:
124         (A) LENGTH: 36 base pairs
125         (B) TYPE: nucleic acid
126         (C) STRANDEDNESS: both
127         (D) TOPOLOGY: both
129     (ii) MOLECULE TYPE: cDNA
134     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
136 GGACCCACTG TCTTTACCGC GGCCTCCTCA AGCACC      36
138 (2) INFORMATION FOR SEQ ID NO: 6:
140     (i) SEQUENCE CHARACTERISTICS:
141         (A) LENGTH: 35 base pairs
142         (B) TYPE: nucleic acid
143         (C) STRANDEDNESS: both
144         (D) TOPOLOGY: both
146     (ii) MOLECULE TYPE: cDNA
151     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
153 CACACACACC CGGGTTAATA CGCTTGGAAG GTGGC      35
155 (2) INFORMATION FOR SEQ ID NO: 7:
157     (i) SEQUENCE CHARACTERISTICS:
158         (A) LENGTH: 31 base pairs
159         (B) TYPE: nucleic acid
160         (C) STRANDEDNESS: both
161         (D) TOPOLOGY: both

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163      (ii) MOLECULE TYPE: cDNA
168      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:
170 CACACACATC ATGACTGTTG CGCTACATCT G      31
172 (2) INFORMATION FOR SEQ ID NO: 8:
174      (i) SEQUENCE CHARACTERISTICS:
175          (A) LENGTH: 29 base pairs
176          (B) TYPE: nucleic acid
177          (C) STRANDEDNESS: both
178          (D) TOPOLOGY: both
180      (ii) MOLECULE TYPE: cDNA
185      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:
187 CACACACAGG TACCCTCTCG AAAAGTTAA      29
189 (2) INFORMATION FOR SEQ ID NO: 9:
191      (i) SEQUENCE CHARACTERISTICS:
192          (A) LENGTH: 48 base pairs
193          (B) TYPE: nucleic acid
194          (C) STRANDEDNESS: both
195          (D) TOPOLOGY: both
197      (ii) MOLECULE TYPE: cDNA
202      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:
204 CACACACAGA GGAATTCAGT GATGGTGATG GTGATGTGCA AAAAGAGG      48
206 (2) INFORMATION FOR SEQ ID NO: 10:
208      (i) SEQUENCE CHARACTERISTICS:
209          (A) LENGTH: 41 base pairs
210          (B) TYPE: nucleic acid
211          (C) STRANDEDNESS: both
212          (D) TOPOLOGY: both
214      (ii) MOLECULE TYPE: cDNA
219      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:
221 ACTGGAATTC ATGCCAATCC ATCACCATCA CCATCACCCG T      41
223 (2) INFORMATION FOR SEQ ID NO: 11:
225      (i) SEQUENCE CHARACTERISTICS:
226          (A) LENGTH: 41 base pairs
227          (B) TYPE: nucleic acid
228          (C) STRANDEDNESS: both
229          (D) TOPOLOGY: both
231      (ii) MOLECULE TYPE: cDNA
236      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:
238 ACGTGTGCGAC CATATGGATG ACTAGGTGAA ACGGGTGATG G      41
240 (2) INFORMATION FOR SEQ ID NO: 12:
242      (i) SEQUENCE CHARACTERISTICS:
243          (A) LENGTH: 71 base pairs
244          (B) TYPE: nucleic acid
245          (C) STRANDEDNESS: both
246          (D) TOPOLOGY: both
248      (ii) MOLECULE TYPE: cDNA
253      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:
255 ACTGGAATTC ATGCCAATCC ATCACCATCA CCATCACCCG TTTCACCTAG TCATCCATAT      60
257 GGTCGACACG T      71

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260 (2) INFORMATION FOR SEQ ID NO: 13:
262     (i) SEQUENCE CHARACTERISTICS:
263         (A) LENGTH: 48 base pairs
264         (B) TYPE: nucleic acid
265         (C) STRANDEDNESS: both
266         (D) TOPOLOGY: both
268     (ii) MOLECULE TYPE: cDNA
273     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:
275 GACTAGTTCT AGATCGCGAG CGGCCGCCCA TTAAGTCTCG TTGGCAGC      48
277 (2) INFORMATION FOR SEQ ID NO: 14:
279     (i) SEQUENCE CHARACTERISTICS:
280         (A) LENGTH: 16 base pairs
281         (B) TYPE: nucleic acid
282         (C) STRANDEDNESS: both
283         (D) TOPOLOGY: both
285     (ii) MOLECULE TYPE: cDNA
290     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:
292 TCGACCCACG CGTCCG      16
294 (2) INFORMATION FOR SEQ ID NO: 15:
296     (i) SEQUENCE CHARACTERISTICS:
297         (A) LENGTH: 12 base pairs
298         (B) TYPE: nucleic acid
299         (C) STRANDEDNESS: both
300         (D) TOPOLOGY: both
302     (ii) MOLECULE TYPE: cDNA
307     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 15:
309 CGGACGCGTG GG      12
311 (2) INFORMATION FOR SEQ ID NO: 16:
313     (i) SEQUENCE CHARACTERISTICS:
314         (A) LENGTH: 42 base pairs
315         (B) TYPE: nucleic acid
316         (C) STRANDEDNESS: both
317         (D) TOPOLOGY: both
319     (ii) MOLECULE TYPE: cDNA
324     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 16:
326 AUGGAGAUCU CUGAATTCAT GACTGTTGCG CTACATCTGG CT      42
328 (2) INFORMATION FOR SEQ ID NO: 17:
330     (i) SEQUENCE CHARACTERISTICS:
331         (A) LENGTH: 32 base pairs
332         (B) TYPE: nucleic acid
333         (C) STRANDEDNESS: both
334         (D) TOPOLOGY: both
336     (ii) MOLECULE TYPE: cDNA
341     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 17:
343 ATTATTCATA TGACTGTTGC GCTACATCTG GC      32
345 (2) INFORMATION FOR SEQ ID NO: 18:
347     (i) SEQUENCE CHARACTERISTICS:
348         (A) LENGTH: 24 base pairs
349         (B) TYPE: nucleic acid

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350         (C) STRANDEDNESS: both
351         (D) TOPOLOGY: both
353         (ii) MOLECULE TYPE: cDNA
358         (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 18:
360 TACGATCTCT CTCCAGGCCA TTTT                                     24
362 (2) INFORMATION FOR SEQ ID NO: 19:
364         (i) SEQUENCE CHARACTERISTICS:
365             (A) LENGTH: 24 base pairs
366             (B) TYPE: nucleic acid
367             (C) STRANDEDNESS: both
368             (D) TOPOLOGY: both
370         (ii) MOLECULE TYPE: cDNA
375         (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19:
377 ACTCGAGCAG CCCGGAACC TTG                                         24
379 (2) INFORMATION FOR SEQ ID NO: 20:
381         (i) SEQUENCE CHARACTERISTICS:
382             (A) LENGTH: 48 base pairs
383             (B) TYPE: nucleic acid
384             (C) STRANDEDNESS: both
385             (D) TOPOLOGY: both
387         (ii) MOLECULE TYPE: cDNA
392         (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:
394 ATTACCCGGG AGGATATCAT ATGTTAGCGA TGACAATGGA ACATAAAG          48
396 (2) INFORMATION FOR SEQ ID NO: 21:
398         (i) SEQUENCE CHARACTERISTICS:
399             (A) LENGTH: 36 base pairs
400             (B) TYPE: nucleic acid
401             (C) STRANDEDNESS: both
402             (D) TOPOLOGY: both
404         (ii) MOLECULE TYPE: cDNA
409         (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 21:
411 ATATGTCGAC TCACAGTGGC CCTCCCTATA AATTTG                         36
413 (2) INFORMATION FOR SEQ ID NO: 22:
415         (i) SEQUENCE CHARACTERISTICS:
416             (A) LENGTH: 35 base pairs
417             (B) TYPE: nucleic acid
418             (C) STRANDEDNESS: both
419             (D) TOPOLOGY: both
421         (ii) MOLECULE TYPE: cDNA
426         (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 22:
428 TATTAGGATC CCATGACTGT TGCGCTACAT CTGGC                         35
430 (2) INFORMATION FOR SEQ ID NO: 23:
432         (i) SEQUENCE CHARACTERISTICS:
433             (A) LENGTH: 29 base pairs
434             (B) TYPE: nucleic acid
435             (C) STRANDEDNESS: both
436             (D) TOPOLOGY: both
438         (ii) MOLECULE TYPE: cDNA
442         (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 23:

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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/245,025

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Input Set : N:\Crf3\RULE60\09245025.txt

Output Set: N:\CRF3\08132001\I245025.raw

L:30 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]

L:31 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]